

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source

Date Processed by STIC:

10/697,720 TFWO 7/6/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER

VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
  U.S. Patent and Trademark Office, 220-20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04



**IFWO** 

RAW SEQUENCE LISTING

DATE: 07/06/2004

TIME: 11:03:58 PATENT APPLICATION: US/10/697,720

Input Set : A:\5398-027-27CONT.ST25.txt Output Set: N:\CRF4\07062004\J697720.raw

- 3 <110 > APPLICANT: Cohen, Stanley N.
- Li, Limin
- 6 <120> TITLE OF INVENTION: Mammalian Tumor Susceptibility Genes and Their Uses
- 8 <130> FILE REFERENCE: 5398-027-27CONT
- 10 <140> CURRENT APPLICATION NUMBER: US 10/697,720
- 11 <141> CURRENT FILING DATE: 2003-10-29
- 13 <150> PRIOR APPLICATION NUMBER: US 09/804,690
- 14 <151> PRIOR FILING DATE: 2001-03-12
- 16 <150> PRIOR APPLICATION NUMBER: US 09/146,187
- 17 <151> PRIOR FILING DATE: 1998-09-01
- 19 <150> PRIOR APPLICATION NUMBER: US 08/977,818
- 20 <151> PRIOR FILING DATE: 1997-11-25
- 22 <150> PRIOR APPLICATION NUMBER: US 08/670,274
- 23 <151> PRIOR FILING DATE: 1996-06-13
- 25 <150> PRIOR APPLICATION NUMBER: US 08/585,758
- 26 <151> PRIOR FILING DATE: 1996-01-16
- 28 <150> PRIOR APPLICATION NUMBER: US 60/006,856
- 29 <151> PRIOR FILING DATE: 1995-11-16
- 31 <160> NUMBER OF SEQ ID NOS: 20
- 33 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply Corrected Diskette Needed

ERRORED SEQUENCES

326 <210> SEQ ID NO: 7

327 <211> LENGTH: \_39

328 <212> TYPE: (RNA)

329 <213> ORGANISM: Artificial Sequence

331 <220> FEATURE:

332 <223> OTHER INFORMATION: primer

334 <400> SEQUENCE: 7

E--> 335 caucaucauc augaggbggc (Babgagbab) Pholificcag

337 <210> SEQ ID NO: 8

338 <211> LENGTH: 39

339 <212> TYPE: (RNA)

340 <213> ORGANISM: Artificial Sequence

342 <220> FEATURE:

343 <223> OTHER INFORMATION: primer

345 <400> SEQUENCE: 8

E--> 346 cuacuacuac uacacdttt gagcaagt cagcd ggtt

39

"És" are allowed = in a RNA sequence.

SAMEERROR

39

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/697,720

DATE: 07/06/2004

TIME: 11:03:59

Input Set : A:\5398-027-27CONT.ST25.txt
Output Set: N:\CRF4\07062004\J697720.raw

L:43 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:46 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:335 M:321 E: (1) "t" not allowed in RNA Sequence, NUMBER OF INVALID 't' KEYS:10
L:346 M:321 E: (1) "t" not allowed in RNA Sequence, NUMBER OF INVALID 't' KEYS:9
L:479 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:483 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0